SEQUENCE LISTING

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Mark J. Graham
Pam Nero
Edward Wancewicz

<120> ANTISENSE MODULATION OF CHOLESTERYL ESTER TRANSFER PROTEIN EXPRESSION

<130> ISPH-0596

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cctgataacc atg ctg gct gcc aca gtc ctg acc ctg gcc ctg ctg ggc 169
Met Leu Ala Ala Thr Val Leu Thr Leu Ala Leu Leu Gly
1 5 10

aat gcc cat gcc tgc tcc aaa ggc acc tcg cac gag gca ggc atc gtg 217
Asn Ala His Ala Cys Ser Lys Gly Thr Ser His Glu Ala Gly Ile Val

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tgc cgc atc acc aag cct gcc ctc ctg gtg ttg aac cac gag act gcc   265
Cys Arg Ile Thr Lys Pro Ala Leu Leu Val Leu Asn His Glu Thr Ala
 30                  35                  40                  45

aag gtg atc cag acc gcc ttc cag cga gcc agc tac cca gat atc acg   313
Lys Val Ile Gln Thr Ala Phe Gln Arg Ala Ser Tyr Pro Asp Ile Thr
                 50                  55                  60

ggc gag aag gcc atg atg ctc ctt ggc caa gtc aag tat ggg ttg cac   361
Gly Glu Lys Ala Met Met Leu Leu Gly Gln Val Lys Tyr Gly Leu His
             65                  70                  75

aac atc cag atc agc cac ttg tcc atc gcc agc agc cag gtg gag ctg   409
Asn Ile Gln Ile Ser His Leu Ser Ile Ala Ser Ser Gln Val Glu Leu
         80                  85                  90

gtg gaa gcc aag tcc att gat gtc tcc att cag aac gtg tct gtg gtc   457
Val Glu Ala Lys Ser Ile Asp Val Ser Ile Gln Asn Val Ser Val Val
     95                  100                 105

ttc aag ggg acc ctg aag tat ggc tac acc act gcc tgg tgg ctg ggt   505
Phe Lys Gly Thr Leu Lys Tyr Gly Tyr Thr Thr Ala Trp Trp Leu Gly
110                 115                 120                 125

att gat cag tcc att gac ttc gag atc gac tct gcc att gac ctc cag   553
Ile Asp Gln Ser Ile Asp Phe Glu Ile Asp Ser Ala Ile Asp Leu Gln
                130                 135                 140

atc aac aca cag ctg acc tgt gac tct ggt aga gtg cgg acc gat gcc   601
Ile Asn Thr Gln Leu Thr Cys Asp Ser Gly Arg Val Arg Thr Asp Ala
            145                 150                 155

cct gac tgc tac ctg tct ttc cat aag ctg ctc ctg cat ctc caa ggg   649
Pro Asp Cys Tyr Leu Ser Phe His Lys Leu Leu Leu His Leu Gln Gly
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gag cga gag cct ggg tgg atc aag cag ctg ttc aca aat ttc atc tcc   697
Glu Arg Glu Pro Gly Trp Ile Lys Gln Leu Phe Thr Asn Phe Ile Ser
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ttc acc ctg aag ctg gtc ctg aag gga cag atc tgc aaa gag atc aac   745
Phe Thr Leu Lys Leu Val Leu Lys Gly Gln Ile Cys Lys Glu Ile Asn
190                 195                 200                 205

gtc atc tct aac atc atg gcc gat ttt gtc cag aca agg gct gcc agc   793
Val Ile Ser Asn Ile Met Ala Asp Phe Val Gln Thr Arg Ala Ala Ser
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atc ctt tca gat gga gac att ggg gtg gac att tcc ctg aca ggt gat   841
Ile Leu Ser Asp Gly Asp Ile Gly Val Asp Ile Ser Leu Thr Gly Asp
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ccc gtc atc aca gcc tcc tac ctg gag tcc cat cac aag ggt cat ttc   889
Pro Val Ile Thr Ala Ser Tyr Leu Glu Ser His His Lys Gly His Phe
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Ile Tyr Lys Asn Val Ser Glu Asp Leu Pro Leu Pro Thr Phe Ser Pro
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aca ctg ctg ggg gac tcc cgc atg ctg tac ttc tgg ttc tct gag cga   985
Thr Leu Leu Gly Asp Ser Arg Met Leu Tyr Phe Trp Phe Ser Glu Arg
270                 275                 280                 285

gtc ttc cac tcg ctg gcc aag gta gct ttc cag gat ggc cgc ctc atg   1033
Val Phe His Ser Leu Ala Lys Val Ala Phe Gln Asp Gly Arg Leu Met
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ctc agc ctg atg gga gac gag ttc aag gca gtg ctg gag acc tgg ggc   1081
Leu Ser Leu Met Gly Asp Glu Phe Lys Ala Val Leu Glu Thr Trp Gly
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ttc aac acc aac cag gaa atc ttc caa gag gtt gtc ggc ggc ttc ccc   1129
Phe Asn Thr Asn Gln Glu Ile Phe Gln Glu Val Val Gly Gly Phe Pro
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agc cag gcc caa gtc acc gtc cac tgc ctc aag atg ccc aag atc tcc   1177
Ser Gln Ala Gln Val Thr Val His Cys Leu Lys Met Pro Lys Ile Ser
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tgc caa aac aag gga gtc gtg gtc aat tct tca gtg atg gtg aaa ttc   1225
Cys Gln Asn Lys Gly Val Val Val Asn Ser Ser Val Met Val Lys Phe
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ctc ttt cca cgc cca gac cag caa cat tct gta gct tac aca ttt gaa   1273
Leu Phe Pro Arg Pro Asp Gln Gln His Ser Val Ala Tyr Thr Phe Glu
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gag gat atc gtg act acc gtc cag gcc tcc tat tct aag aaa aag ctc   1321
Glu Asp Ile Val Thr Thr Val Gln Ala Ser Tyr Ser Lys Lys Lys Leu
            385                 390                 395

ttc tta agc ctc ttg gat ttc cag att aca cca aag act gtt tcc aac   1369
Phe Leu Ser Leu Leu Asp Phe Gln Ile Thr Pro Lys Thr Val Ser Asn
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Leu Thr Glu Ser Ser Ser Glu Ser Ile Gln Ser Phe Leu Gln Ser Met
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atc acc gct gtg ggc atc cct gag gtc atg tct cgg ctc gag gta gtg   1465
Ile Thr Ala Val Gly Ile Pro Glu Val Met Ser Arg Leu Glu Val Val
430                 435                 440                 445

ttt aca gcc ctc atg aac agc aaa ggc gtg agc ctc ttc gac atc atc   1513
Phe Thr Ala Leu Met Asn Ser Lys Gly Val Ser Leu Phe Asp Ile Ile
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aac cct gag att atc act cga gat ggc ttc ctg ctg ctg cag atg gac   1561
Asn Pro Glu Ile Ile Thr Arg Asp Gly Phe Leu Leu Leu Gln Met Asp
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gctcacagct ggaaccctgg tgtctcctcc agcgtggtgg aagttgggtt aggagtacgg 1722
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